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See page 6



1600

RAW SEQUENCE LISTING

DATE: 06/03/2002

PATENT APPLICATION: US/09/492,764B

TIME: 13:31:22

Input Set : A:\10873-008.txt

Output Set: N:\CRF3\06032002\I492764B.raw

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JUN 12 2002

TECH CENTER 1600/2900

5 <110> APPLICANT: Jove, Richard
 7 Hamilton, Andrew
 9 Gilbert, Richard
 11 Dalton, William
 13 Sebti, Said
 15 Yu, Hua
 17 Heller, Richard
 19 Jaroszeski, Mark
 23 <120> TITLE OF INVENTION: INHIBITION OF STAT3 SIGNAL TRANSDUCTION FOR HUMAN CANCER
 24 THERAPY
 28 <130> FILE REFERENCE: 10873-008-999
 32 <140> CURRENT APPLICATION NUMBER: 09/492,764B
 34 <141> CURRENT FILING DATE: 2000-01-27
 38 <150> PRIOR APPLICATION NUMBER: 60/117,600
 40 <151> PRIOR FILING DATE: 1999-01-27
 44 <160> NUMBER OF SEQ ID NOS: 38
 48 <170> SOFTWARE: PatentIn version 3.0
 52 <210> SEQ ID NO: 1
 54 <211> LENGTH: 24
 56 <212> TYPE: DNA
 58 <213> ORGANISM: Homo sapiens
 62 <400> SEQUENCE: 1
 63 agcttcattt cccgtaaata ccta 24
 66 <210> SEQ ID NO: 2
 68 <211> LENGTH: 24
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 72 <213> ORGANISM: Homo sapiens
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 80 <210> SEQ ID NO: 3
 82 <211> LENGTH: 22
 84 <212> TYPE: DNA
 86 <213> ORGANISM: Homo sapiens
 90 <400> SEQUENCE: 3
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 94 <210> SEQ ID NO: 4
 96 <211> LENGTH: 25
 98 <212> TYPE: DNA
 100 <213> ORGANISM: Homo sapiens
 104 <400> SEQUENCE: 4
 105 cgacgacttc tcccgcgcgt accgc 25
 108 <210> SEQ ID NO: 5
 110 <211> LENGTH: 25

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112 <212> TYPE: DNA
114 <213> ORGANISM: Homo sapiens
118 <400> SEQUENCE: 5
119 ccgcatgctg gggccgtaca gttcc 25
122 <210> SEQ ID NO: 6
124 <211> LENGTH: 20
126 <212> TYPE: DNA
128 <213> ORGANISM: Homo sapiens
132 <400> SEQUENCE: 6
133 cgggcattca gtgacctgac 20
136 <210> SEQ ID NO: 7
138 <211> LENGTH: 20
140 <212> TYPE: DNA
142 <213> ORGANISM: Homo sapiens
146 <400> SEQUENCE: 7
147 tcaggaacca gcggttgaag 20
150 <210> SEQ ID NO: 8
152 <211> LENGTH: 20
154 <212> TYPE: DNA
156 <213> ORGANISM: Homo sapiens
160 <400> SEQUENCE: 8
161 ccactgaact tctgattcgc 20
164 <210> SEQ ID NO: 9
166 <211> LENGTH: 20
168 <212> TYPE: DNA
170 <213> ORGANISM: Homo sapiens
174 <400> SEQUENCE: 9
175 gcgtgctagc tggatgtcctt 20
178 <210> SEQ ID NO: 10
180 <211> LENGTH: 9
182 <212> TYPE: DNA
184 <213> ORGANISM: Homo sapiens
188 <400> SEQUENCE: 10
189 ttcggagaa 9
192 <210> SEQ ID NO: 11
194 <211> LENGTH: 9
196 <212> TYPE: DNA
198 <213> ORGANISM: Homo sapiens
202 <400> SEQUENCE: 11
203 tgaggataa 9
206 <210> SEQ ID NO: 12
208 <211> LENGTH: 12
210 <212> TYPE: PRT
212 <213> ORGANISM: Homo sapiens
216 <220> FEATURE:
218 <221> NAME/KEY: misc_feature
220 <223> OTHER INFORMATION: Xaa = Ser or Pro
224 <400> SEQUENCE: 12
W--> 226 His Tyr Xaa Pro Ile Leu Val Tyr Gln Pro Ser Trp

```

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229 <210> SEQ ID NO: 13
231 <211> LENGTH: 12
233 <212> TYPE: PRT
235 <213> ORGANISM: Homo sapiens
239 <400> SEQUENCE: 13
241 Gln Asp Val His Leu Thr Gln Gln Ser Arg Tyr Thr
242 1          5          10
244 <210> SEQ ID NO: 14
246 <211> LENGTH: 12
248 <212> TYPE: PRT
250 <213> ORGANISM: Homo sapiens
254 <400> SEQUENCE: 14
256 Ser His Pro Trp Asn Ala Gln Arg Glu Leu Ser Val
257 1          5          10
259 <210> SEQ ID NO: 15
261 <211> LENGTH: 12
263 <212> TYPE: PRT
265 <213> ORGANISM: Homo sapiens
269 <400> SEQUENCE: 15
271 Tyr Pro Ala Pro Gln Pro Leu Val Thr Lys Thr Ser
272 1          5          10
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276 <211> LENGTH: 12
278 <212> TYPE: PRT
280 <213> ORGANISM: Homo sapiens
284 <400> SEQUENCE: 16
286 Phe Ser Tyr Pro Leu Thr Arg Ala Pro Leu Asn Met
287 1          5          10
289 <210> SEQ ID NO: 17
291 <211> LENGTH: 7
293 <212> TYPE: PRT
295 <213> ORGANISM: Homo sapiens
299 <400> SEQUENCE: 17
301 His Ala Ile Tyr Pro Arg Asn
302 1          5
304 <210> SEQ ID NO: 18
306 <211> LENGTH: 7
308 <212> TYPE: PRT
310 <213> ORGANISM: Homo sapiens
314 <400> SEQUENCE: 18
316 Ala Ser Thr Leu Pro Lys Ala
317 1          5
319 <210> SEQ ID NO: 19
321 <211> LENGTH: 7
323 <212> TYPE: PRT
325 <213> ORGANISM: Homo sapiens
329 <400> SEQUENCE: 19
331 Ile Gln Ser Pro His Phe Phe
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334 <210> SEQ ID NO: 20
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344 <220> FEATURE:
346 <221> NAME/KEY: misc_feature
348 <223> OTHER INFORMATION: X = PHOSPHOTYROSINE
352 <400> SEQUENCE: 20
W--> 354 Pro Xaa Leu Lys Thr Lys
355 1      5
357 <210> SEQ ID NO: 21
359 <211> LENGTH: 6
361 <212> TYPE: PRT
363 <213> ORGANISM: Homo sapiens
367 <400> SEQUENCE: 21
369 Pro Tyr Leu Lys Thr Lys
370 1      5
372 <210> SEQ ID NO: 22
374 <211> LENGTH: 6
376 <212> TYPE: PRT
378 <213> ORGANISM: Homo sapiens
382 <220> FEATURE:
384 <221> NAME/KEY: misc_feature
386 <223> OTHER INFORMATION: X= PHOSPHOTYROSINE
390 <400> SEQUENCE: 22
W--> 392 Ala Xaa Leu Lys Thr Lys
393 1      5
395 <210> SEQ ID NO: 23
397 <211> LENGTH: 6
399 <212> TYPE: PRT
401 <213> ORGANISM: Homo sapiens
405 <220> FEATURE:
407 <221> NAME/KEY: misc_feature
409 <223> OTHER INFORMATION: X = PHOSPHOTYROSINE
413 <400> SEQUENCE: 23
W--> 415 Pro Xaa Ala Lys Thr Lys
416 1      5
418 <210> SEQ ID NO: 24
420 <211> LENGTH: 6
422 <212> TYPE: PRT
424 <213> ORGANISM: Homo sapiens
428 <220> FEATURE:
430 <221> NAME/KEY: misc_feature
432 <223> OTHER INFORMATION: X = PHOSPHOTYROSINE
436 <400> SEQUENCE: 24
W--> 438 Pro Xaa Leu Ala Thr Lys
439 1      5
441 <210> SEQ ID NO: 25
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Input Set : A:\10873-008.txt

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443 <211> LENGTH: 6
445 <212> TYPE: PRT
447 <213> ORGANISM: Homo sapiens
451 <220> FEATURE:
453 <221> NAME/KEY: misc_feature
455 <223> OTHER INFORMATION: X = PHOSPHOTYROSINE
459 <400> SEQUENCE: 25
W--> 461 Pro Xaa Leu Lys Ala Lys
462 1 5
464 <210> SEQ ID NO: 26
466 <211> LENGTH: 6
468 <212> TYPE: PRT
470 <213> ORGANISM: Homo sapiens
474 <400> SEQUENCE: 26
476 Pro Tyr Leu Lys Thr Ala
477 1 5
479 <210> SEQ ID NO: 27
481 <211> LENGTH: 4
483 <212> TYPE: PRT
485 <213> ORGANISM: Homo sapiens
489 <220> FEATURE:
491 <221> NAME/KEY: misc_feature
493 <223> OTHER INFORMATION: X = PHOSPHOTYROSINE
497 <400> SEQUENCE: 27
W--> 499 Pro Xaa Leu Lys
500 1
502 <210> SEQ ID NO: 28
504 <211> LENGTH: 4
506 <212> TYPE: PRT
508 <213> ORGANISM: Homo sapiens
512 <220> FEATURE:
514 <221> NAME/KEY: misc_feature
516 <223> OTHER INFORMATION: X = PHOSPHOTYROSINE
520 <400> SEQUENCE: 28
W--> 522 Pro Xaa Phe Lys
523 1
525 <210> SEQ ID NO: 29
527 <211> LENGTH: 3
529 <212> TYPE: PRT
531 <213> ORGANISM: Homo sapiens
535 <220> FEATURE:
537 <221> NAME/KEY: misc_feature
539 <223> OTHER INFORMATION: X = PHOSPHOTYROSINE
543 <400> SEQUENCE: 29
W--> 545 Xaa Leu Lys
546 1
548 <210> SEQ ID NO: 30
550 <211> LENGTH: 4
552 <212> TYPE: PRT
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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/492,764B

DATE: 06/03/2002
TIME: 13:31:23

Input Set : A:\10873-008.txt
Output Set: N:\CRF3\06032002\I492764B.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:12; Xaa Pos. 3
Seq#:20; Xaa Pos. 2
Seq#:22; Xaa Pos. 2
Seq#:23; Xaa Pos. 2
Seq#:24; Xaa Pos. 2
Seq#:25; Xaa Pos. 2
Seq#:27; Xaa Pos. 2
Seq#:28; Xaa Pos. 2
Seq#:29; Xaa Pos. 1
Seq#:30; Xaa Pos. 2
Seq#:32; Xaa Pos. 2
Seq#:34; Xaa Pos. 2
Seq#:35; Xaa Pos. 2
Seq#:36; Xaa Pos. 2
Seq#:37; Xaa Pos. 2
Seq#:38; Xaa Pos. 2

VARIABLE LOCATION SUMMARY
PATENT APPLICATION: US/09/492,764B

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Input Set : A:\10873-008.txt
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Use of n's or Xaa's(NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing.

Use of <220> to <223> is MANDATORY if n's or Xaa's are present.

in <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

Seq#:12; Xaa Pos. 3
Seq#:20; Xaa Pos. 2
Seq#:22; Xaa Pos. 2
Seq#:23; Xaa Pos. 2
Seq#:24; Xaa Pos. 2
Seq#:25; Xaa Pos. 2
Seq#:27; Xaa Pos. 2
Seq#:28; Xaa Pos. 2
Seq#:29; Xaa Pos. 1
Seq#:30; Xaa Pos. 2
Seq#:32; Xaa Pos. 2
Seq#:34; Xaa Pos. 2
Seq#:35; Xaa Pos. 2
Seq#:36; Xaa Pos. 2
Seq#:37; Xaa Pos. 2
Seq#:38; Xaa Pos. 2

VERIFICATION SUMMARY

DATE: 06/03/2002

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Input Set : A:\10873-008.txt

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L:226 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:12
L:226 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0
L:354 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:20
L:354 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0
L:392 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:22
L:392 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0
L:415 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:23
L:415 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0
L:438 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:24
L:438 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0
L:461 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:25
L:461 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0
L:499 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:27
L:499 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0
L:522 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:28
L:522 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:0
L:545 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:29
L:545 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29 after pos.:0
L:568 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:30
L:568 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:0
L:606 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:32
L:606 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:0
L:644 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:34
L:644 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 after pos.:0
L:667 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:35
L:667 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:0
L:690 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:36
L:690 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:0
L:713 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:37
L:713 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:0
L:736 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:38
L:736 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38 after pos.:0